Project Title	Funding	Institution	
Whole-exome sequencing to identify causative genes for autism	\$134,203	ROCKEFELLER UNIVERSITY	
Validation of candidate ASD genes by targeted sequencing with molecular inversion probes	\$101,258	The Regents of the University of California, San Francisco (Contracts & Grants)	
Use of High-throughput Splicing Assays to Prioritize Autism Gene Candidates	\$62,500	Brown University	
University of Washington Clinical Site Network Pilot for the National Autism Cohort	\$37,500	University of Washington	
University of North Carolina Clinical Site Network Pilot for the National Autism Cohort	\$24,750	University of North Carolina	
Thompson Center Clinical Site Network Pilot for the National Autism Cohort	\$37,500	The Curators of the University of Missouri	
The Future of Genomics Medicine in Patient Care: Contributions from CHOP	\$906,296	Children's Hospital of Philadelphia	
Structural Variation and the Genetic Architecture of Autism	\$0	University of Washington	
Simons Simplex Collection support grant	\$10,000	University of Washington	
Simons Simplex Collection support grant	\$9,159	University of Illinois at Chicago	
Simons Simplex Collection support grant	\$8,912	Vanderbilt University	
Simons Simplex Collection support grant	\$13,200	University of California, Los Angeles	
Simons Simplex Collection support grant	\$5,983	Baylor College of Medicine	
Simons Simplex Collection support grant	\$1,831	Weill Cornell Medical College	
Simons Simplex Collection support grant	\$10,000	University of Missouri	
Simons Simplex Collection support grant	\$8,800	Emory University	
Simons Simplex Collection support grant	\$10,000	McGill University Health Centre- Montreal Children's Hospital	
Simons Simplex Collection support grant	\$10,000	Yale University	
Sequencing Female-enriched Multiplex Autism Families (FEMFs)	\$0	Johns Hopkins University	
Sequence-based discovery of genes with pleiotropic effects across diagnostic boundaries and throughout the lifespan	\$14,998	Massachusetts General Hospital	
Role of Selfish Spermatogonial Selection in Neurocognitive Disorders	\$59,995	University of Oxford	
Pieces of the Puzzle: Uncovering the Genetics of Autism	\$1,699,790	Broad Institute, Inc.	
Novel Statistical methods for DNA Sequencing Data, and applications to Autism.	\$318,575	Columbia University	
Next Generation Gene Discovery in Familial Autism	\$644,823	University of Washington	
New York Genome Center, Inc.	\$2,210,000	New York Genome Center, Inc.	
Mutations in noncoding DNA and the missing heritability of autism	\$244,030	University of California, San Diego	
MSSNG	\$5,802,895	Autism Speaks (AS)	
Investigating the role of somatic mutations in autism spectrum disorders	\$263,892	OREGON HEALTH & SCIENCE UNIVERSITY	
Integrating the genomics of Autism Spectrum Disorders(ASD) in consanguineous and "idiopathic" families	\$665,939	Yale University	
Integrating large scale whole exome data with whole genome data	\$0	ICAHN SCHOOL OF MEDICINE AT MOUNT SINAI	
Illumina, Inc.	\$0	Illumina, Inc.	

Project Title	Funding	Institution	
Identifying Patterns of Genetic Variants Conferring Risk for Neurodevelopmental Disorders	\$29,987	Pennsylvania State University	
Identification and analysis of functional networks perturbed in autism	\$250,000	Columbia University	
High-throughput Screening of Novel Trinucleotide Repeat Expansion in Autism Spectrum Disorders	\$15,000	The Hospital for Sick Children	
Genomic profiling of autism families using whole-genome sequencing	\$0	Institut Pasteur	
Genomic influences on development and outcomes in infants at risk for autism	\$0	University of Alberta	
Genome-wide analysis of cis-regulatory elements in autism	\$62,500	Washington University in St. Louis	
Genome Sequencing pilot of Simons Simplex Collection	\$0	University of Washington	
Genetic Epidemiology of Complex Traits	\$747,204	National Institutes of Health	
Genetic basis of phenotypic variability in 16p11.2 deletion or duplication	\$285,856	University of Washington	
Genetic basis of autism	\$4,000,000	Cold Spring Harbor Laboratory	
FUNCTIONAL DISSECTION OF CNVS IN NEURODEVELOPMENTAL TRAITS	\$366,666	Duke University	
Extending ASD risk locus discovery to the non-coding genome - Project 2	\$0	Yale University	
Extending ASD risk locus discovery to the non-coding genome - Project 1	\$0	The Trustees of Columbia University in the City of New York	
Extending ASD risk locus discovery to the non-coding genome - Core	\$0	The Regents of the University of California, San Francisco (Contracts & Grants)	
Elucidating pathogenic mutations disrupting RNA regulation in autism	\$225,000	Columbia University	
Dosage effects of DUF1220 gene subtype CON1 in autism	\$125,000	University of Colorado, Denver	
Discovery of regulatory variants underlying pediatric neurological disease	\$0	HudsonAlpha Institute for Biotechnology	
DEVELOPING NEW STATISICAL METHODS TO DETECT RARE VARIANTS INVOLVED IN NEUROPSYCHIATRIC DISORDERS	\$433,800	National Institutes of Health	
Cryptic Genetic Causes of Autism	\$141,719	Massachusetts General Hospital	
Computational tools to analyze SNP data from patients with mental illness	\$572,792	PARTEK, INC.	
Complex Genetic Architecture of Chromosomal Aberrations in Autism	\$248,999	Massachusetts General Hospital	
Combining WGS from Utah high-risk pedigrees and SSC families	\$0	University of Utah	
Autism Spectrum Disorders: Genomes to Outcomes	\$0	Hospital for Sick Children	
Autism genetics: homozygosity mapping and functional validation	\$765,736	CHILDREN'S HOSPITAL CORPORATION	
ASD Family Biobank Program	\$0	Kaiser Foundation Research Institute	
Applications of novel statistical methods to CNVs in autism and schizophrenia	\$200,000	Columbia University	
Accelerating Autism Genetics via Whole Population Ascertainment in Denmark	\$0	Broad Institute, Inc.	
3/3-Sequencing Autism Spectrum Disorder Extended Pedigrees	\$160,000	University of Pennsylvania	

Project Title	Funding	Institution
2/3 Sequencing Autism Spectrum Disorder Extended Pedigrees	\$231,750	University of Washington
1/3 - Sequencing Autism Spectrum Disorder Extended Pedigrees	\$298,000	UNIVERSITY OF UTAH